



SEQUENCE LISTING

<110> Institut National de la Recherche Agronomique (INRA)

<120> Method of producing double low restorer lines of Brassica napus having a good agronomic value

<130> D21413

<140> US 10/563,277

<141> 2006-01-04

<160> 14

<170> PatentIn version 3.2

<210> 1

<211> 248

<212> DNA

<213> Brassica napus

<220>

<223> PGIo1 marker

<400> 1

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aatcttgctg	tatgaatttg	tgattaaatt	tgtttggttg	tgactctttc	ttcattgttc	180
gttttcgtac	aataaaccga	atgtataatc	ttttacaaa	ctgaattttc	taccgggtct	240
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<210> 2

<211> 979

<212> DNA

<213> Brassica napus

<220>

<223> PGI-UNT R2000 marker

<400> 2

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gtaagtgcct	gtttatttgg	ttgtataaat	ttctcgtcca	tttccgcttg	cttagtgtat	480
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agttgtgtga	ttatacagtt	ttcttgctct	tttgctatgt	ccatcaacac	tagagattcg	900
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 <212> DNA
 <213> Brassica napus

<220>
 <223> PGI-int R2000 marker

<400> 3

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aaaatgtctt	aggcatattc	tttctathtt	atctccctct	taatgatttc	ttcttttttt	540
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ttgcattagt	tgtgtgatta	tacagttttc	ttgtcttttt	gctatgtcca	tcaacactag	780
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 <211> 957
 <212> DNA
 <213> Brassica napus

<220>
 <223> BolJon marker R2000

<400> 4

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gggataacag	tgtagaaaac	aaaccgtctg	taagattttc	tccctgatcc	tctcacttaa	180
ccagtagggc	tttttcacat	tgaagcgc	atctactttg	gtattcactg	aataaaaaaa	240
gaaagctgg	aacatgtgaa	ggatatacaa	gcattgatac	accaagtagt	cacaaactac	300
attataaagg	tcagaccttt	gttcacattc	tggtctccag	gaccaccgct	tctagcaaag	360
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atacaaacaa	aactatgcga	acagatcaaa	actactacag	aacacagttc	tatgacactg	540
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 <212> DNA
 <213> Brassica napus

<220>
 <223> CP418L marker R2000

<400> 5

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ccactacata	gcagagttct	tataacattg	tctgtaaatg	tccaatcaaa	accactacag	180
aacaaagctc	ctataacatt	gtttatacaa	agtttctacta	aatctacaaa	ctttccccgt	240
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gagattagaa	caaactgaaa	cttacgtaga	gtgatttgag	gagtaggctc	gttgccagca	360
gagctagctc	tctcctccgc	ctcatgaagc	atctgttgca	cctgagacaa	ccgtgacgaa	420
actttccgat	caccgccacc	agaattcgac	gccgcgcac	ggaaggatcc	gaatcgggaa	480
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 <223> PGIol U primer

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<210> 7
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 <212> DNA
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<220>
 <223> PGIol L primer

<400> 7
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<210> 11
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<400> 11
 caataaccct aaaagcacct g

21

<210> 12
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<220>
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<400> 12
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21

<210> 13
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<400> 13
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21

<210> 14
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<220>
 <223> pCP418 L primer

<400> 14
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21